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APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO.
09/899,513	07/05/2001	Yi Hu	LEX-0200-USA	9922

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LEXICON GENETICS INCORPORATED  
8800 TECHNOLOGY FOREST PLACE  
THE WOODLANDS, TX 77381-1160

EXAMINER

LI, RUIXIANG

ART UNIT	PAPER NUMBER
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1646

DATE MAILED: 04/23/2003

14

Please find below and/or attached an Office communication concerning this application or proceeding.

**Advisory Action**

Application No.

09/899,513

Applicant(s)

HU ET AL.

Examiner

Ruixiang Li

Art Unit

1646

--The MAILING DATE of this communication appears on the cover sheet with the correspondence address --

THE REPLY FILED 20 March 2003 FAILS TO PLACE THIS APPLICATION IN CONDITION FOR ALLOWANCE. Therefore, further action by the applicant is required to avoid abandonment of this application. A proper reply to a final rejection under 37 CFR 1.113 may only be either: (1) a timely filed amendment which places the application in condition for allowance; (2) a timely filed Notice of Appeal (with appeal fee); or (3) a timely filed Request for Continued Examination (RCE) in compliance with 37 CFR 1.114.

**PERIOD FOR REPLY** [check either a) or b)]

- a) ☒ The period for reply expires 3 months from the mailing date of the final rejection.
- b) ☐ The period for reply expires on: (1) the mailing date of this Advisory Action, or (2) the date set forth in the final rejection, whichever is later. In no event, however, will the statutory period for reply expire later than SIX MONTHS from the mailing date of the final rejection. ONLY CHECK THIS BOX WHEN THE FIRST REPLY WAS FILED WITHIN TWO MONTHS OF THE FINAL REJECTION. See MPEP 706.07(f).

Extensions of time may be obtained under 37 CFR 1.136(a). The date on which the petition under 37 CFR 1.136(a) and the appropriate extension fee have been filed is the date for purposes of determining the period of extension and the corresponding amount of the fee. The appropriate extension fee under 37 CFR 1.17(a) is calculated from: (1) the expiration date of the shortened statutory period for reply originally set in the final Office action; or (2) as set forth in (b) above, if checked. Any reply received by the Office later than three months after the mailing date of the final rejection, even if timely filed, may reduce any earned patent term adjustment. See 37 CFR 1.704(b).

1. ☐ A Notice of Appeal was filed on \_\_\_\_\_. Appellant's Brief must be filed within the period set forth in 37 CFR 1.192(a), or any extension thereof (37 CFR 1.191(d)), to avoid dismissal of the appeal.
2. ☐ The proposed amendment(s) will not be entered because:
- (a) ☐ they raise new issues that would require further consideration and/or search (see NOTE below);
  - (b) ☐ they raise the issue of new matter (see Note below);
  - (c) ☐ they are not deemed to place the application in better form for appeal by materially reducing or simplifying the issues for appeal; and/or
  - (d) ☐ they present additional claims without canceling a corresponding number of finally rejected claims.

NOTE: \_\_\_\_\_

3. ☐ Applicant's reply has overcome the following rejection(s): \_\_\_\_\_.
4. ☐ Newly proposed or amended claim(s) \_\_\_\_\_ would be allowable if submitted in a separate, timely filed amendment canceling the non-allowable claim(s).
5. ☒ The a) ☐ affidavit, b) ☐ exhibit, or c) ☒ request for reconsideration has been considered but does NOT place the application in condition for allowance because: See Continuation Sheet.
6. ☐ The affidavit or exhibit will NOT be considered because it is not directed SOLELY to issues which were newly raised by the Examiner in the final rejection.
7. ☒ For purposes of Appeal, the proposed amendment(s) a) ☐ will not be entered or b) ☒ will be entered and an explanation of how the new or amended claims would be rejected is provided below or appended.

The status of the claim(s) is (or will be) as follows:

Claim(s) allowed: \_\_\_\_\_.

Claim(s) objected to: \_\_\_\_\_.

Claim(s) rejected: 5-7.

Claim(s) withdrawn from consideration: \_\_\_\_\_.

8. ☐ The proposed drawing correction filed on \_\_\_\_\_ is a) ☐ approved or b) ☐ disapproved by the Examiner.
9. ☐ Note the attached Information Disclosure Statement(s) (PTO-1449) Paper No(s). \_\_\_\_\_.
10. ☐ Other: \_\_\_\_\_

Continuation of 5. does NOT place the application in condition for allowance because: the rejection of claims 5-7 under 35 U.S.C. §101 and §112, 1st paragraph remains.

**I. Claim rejection under 35 U.S.C. § 101**

The instant specification fails to satisfy the utility requirement set forth under 35 U.S.C. § 101 for the following reasons, as well as for the reasons set forth in the previous office actions (Paper No. 9 and Paper No. 12).

Applicants argue that the references of Bork and Koonin, Ji, and Yan, do not support the Examiner's position in rejecting claims 5-7 for lack of a patentable utility. This has been fully considered but is not deemed to be persuasive for the following reasons.

Bork and Koonin's conclusion remarks clearly indicates that the potential importance of sequence analysis in extracting functional signal. However, Bork and Koonin do not teach, in any means, that sequence analysis alone can define the biological functions. In fact, Bork and Koonin further teach many proteins are multifunctional, assignment of a single function, which is still common in genome projects, results in loss of information and outright errors (Table 2). As the Examiner stated in the previous office in paper No. 12, while sequence analysis is important, the information provided or "predicted" based upon sequence homology can only be used as guidance in determining functions or activities of a molecule by experiments. Any functions predicted based upon the sequence homology will have to be confirmed ultimately by bench work.


Applicants also argue that an exact quote from Ji completely undermines the question of asserted utility based on sequence homology. The Examiner disagrees. The cited statement simply indicates that a substantial degree of amino acid homology is found between members of a particular subfamily. However, two sequences sharing certain degree homology may not necessarily belong to the same subfamily. In addition, the instant disclosure merely asserts that the polynucleotides of the present invention encode proteins that share sequence similarity with mammalian membrane ligand proteins or transporter proteins (1<sup>st</sup> paragraph of page 2). The disclosure fails to specify a functional protein which the protein of the present invention shares sequence homology with and the degree of homology. Ji also clearly teach there are putative seven transmembrane molecules, which do not appear to be coupled to a G protein. Even if the protein of the present invention were a member of the GPCR family, it would still not provide a patentable utility for the claimed invention because it still requires undue experimentation to define the specific biological function of the present protein or nucleic acid.

Applicants further argue that Yan does not suggest a high level of uncertainty in assigning function based on sequence, and thus does not support the lack of utility. Specifically, applicants argue that the different receptors bound by the two isoforms of ectodysplasin are related and EDA-A2 receptor was correctly identified as a member of the tumor necrosis factor receptor superfamily based upon solely on sequence similarity. The Examiner notes that while the two receptors bound by the two isoforms of ectodysplasin are related, i.e., belonging to the TNFR superfamily, they have different activities (See, e.g., page 524, column 3) and are distinct receptors. Even the title of the paper clearly states that the two receptors bound by the two isoforms are distinct. The Examiner further notes that while the EDA-A2 receptor was initially identified as a member of the TNFR superfamily solely based on sequence similarity, as applicants argued, the biological functions of the receptor was not identified. In fact, Yan et al. performed undue experimentation to define the ligand and biological activities of the receptor. As taught by Yan, the members of the TNFR superfamily are involved in a number of physiological and pathological response by activating a wide variety of intracellular signaling pathways (beginning of page 523). The EDA-A2 receptor (XEDAR) fails to bind many known ligands of the TNF superfamily (1<sup>st</sup> column of page 524). Therefore, even if sequence analysis could assign a given protein to a protein family, the protein still does not have a substantial utility because the biological function or activity is not defined. Determining such a biological function of the protein would require significant further research, as demonstrated by Yan, which is not allowed under 35 U.S.C. § 101.

In addition, applicants continue to argue for the issues of record (Paper No. 11, December 5, 2002). Applicants' arguments were addressed in the previous office action in Paper No. 12.

**II. Claim Rejections Under 35 U. S. C. §112, 1st Paragraph**

Claims 5-7 are rejected under 35 U. S. C. 112, 1st paragraph. Specifically, since the claimed invention is not supported by either a specific, substantial, and credible utility, or a well-established utility, one skilled in the art clearly would not know how to use the claimed invention. The applicants' arguments about the patentable utility of the claimed invention has been fully considered but is not deemed to be persuasive for reason set for the above.

  
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